## SEQUENCE LISTING

<110> University of Illinois at Chicago Sharma, Arun Hoffman, Ronald

<120> HUMAN HEMATOPOIETIC GROWTH REGULATORY GENE AND USES

<130> MBHB: CU08/PPA

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<170> PatentIn version 3.1

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≨213> Homo sapiens

**=** 220>

₹221> CDS

¥222> (1)..(2328)

₹223> Human Hiwi Protein

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Ser Lys Thr Gly Ser Ser Gly Ile Ile Val Arg Leu Ser Thr Asn His

20
25
30

48

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Phe Arg Leu Thr Ser Arg Pro Gln Trp Ala Leu Tyr Gln Tyr His Ile

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Asp Tyr Asn Pro Leu Met Glu Ala Arg Arg Leu Arg Ser Ala Leu Leu

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Phe Gln His Glu Asp Leu Ile Gly Lys Cys His Ala Phe Asp Gly Thr

70 75 80

55

Page 1

60

|     |   |      |  | - |   |   |   |   |   |   |   | gtt<br>Val<br>95  |   | 288 |
|-----|---|------|--|---|---|---|---|---|---|---|---|-------------------|---|-----|
| _   | _ |      |  |   | - |   |   |   |   |   |   | tta<br>Leu        |   | 336 |
|     | _ |      |  |   |   |   | _ | _ | _ |   |   | aat<br>Asn        |   | 384 |
|     |   | <br> |  |   |   | - |   |   |   |   |   | gga<br>Gly        | _ | 432 |
|     |   |      |  | _ |   |   | _ |   |   | _ |   | agg<br>Arg        | _ | 480 |
| gtg |   |      |  |   |   |   |   |   | _ |   | _ | aac<br>Asn<br>175 | _ | 528 |
|     |   |      |  |   | _ |   |   |   |   |   |   | gag<br>Glu        |   | 576 |
|     |   |      |  |   |   |   |   |   |   |   |   | cat<br>His        |   | 624 |
|     |   |      |  |   |   |   |   |   |   | _ | _ | ctt<br>Leu        |   | 672 |
|     |   |      |  |   |   |   |   |   |   | _ |   | gac<br>Asp        | _ | 720 |
|     |   |      |  |   |   |   |   |   |   |   |   | agc<br>Ser<br>255 |   | 768 |
|     |   |      |  |   |   |   |   |   |   |   |   | ttg<br>Leu        |   | 816 |

|            |  | 260 |   |   | 265 |  |       | 270 |                   |          |
|------------|--|-----|---|---|-----|--|-------|-----|-------------------|----------|
|            |  |     |   |   |     |  |       |     | gly<br>aaa        | 864      |
|            |  |     | _ | _ |     |  | _     |     | ctt<br>Leu        | 912      |
|            |  |     |   |   |     |  | <br>_ |     | gac<br>Asp        | 960      |
|            |  |     |   |   |     |  |       |     | gtg<br>Val<br>335 | <br>1008 |
| Arg        |  |     |   |   |     |  |       |     | gag<br>Glu        | 1056     |
|            |  |     |   |   |     |  |       |     | tca<br>Ser        | <br>1104 |
| aga<br>Arg |  |     |   |   |     |  |       |     | ttt<br>Phe        | 1152     |
|            |  |     |   |   |     |  |       |     | gca<br>Ala        | 1200     |
|            |  |     |   |   |     |  |       |     | acg<br>Thr<br>415 | 1248     |
|            |  |     |   |   |     |  |       |     | aaa<br>Lys        | 1296     |
|            |  |     |   |   |     |  |       |     | gaa<br>Glu        | 1344     |

|            |                   |                   |                   |                   | gcc<br>Ala        |                   |                   |                   |                   |                   |                   |                   |                   |                   |                   | 1392 |
|------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|------|
|            |                   |                   |                   |                   | gtt<br>Val<br>470 | _                 | _                 | _                 | _                 |                   | _                 |                   |                   | _                 | _                 | 1440 |
|            |                   |                   |                   |                   | aaa<br>Lys        |                   |                   |                   |                   |                   | _                 | _                 |                   |                   |                   | 1488 |
|            |                   |                   |                   |                   | gcc<br>Ala        |                   |                   |                   |                   |                   |                   |                   |                   | _                 | _                 | 1536 |
| gcc<br>Ala | att<br>Ile        | gct<br>Ala<br>515 | aca<br>Thr        | aag<br>Lys        | att<br>Ile        | gcc<br>Ala        | cta<br>Leu<br>520 | cag<br>Gln        | atg<br>Met        | aac<br>Asn        | tgc<br>Cys        | aag<br>Lys<br>525 | atg<br>Met        | gga<br>Gly        | gga<br>Gly        | 1584 |
| gag        |                   |                   |                   |                   | gac<br>Asp        |                   |                   |                   |                   |                   |                   |                   |                   |                   |                   | 1632 |
| Tle<br>545 | gat<br>Asp        | tgt<br>Cys        | tac<br>Tyr        | cat<br>His        | gac<br>Asp<br>550 | atg<br>Met        | aca<br>Thr        | gct<br>Ala        | gly<br>ggg        | cgg<br>Arg<br>555 | agg<br>Arg        | tca<br>Ser        | atc<br>Ile        | gca<br>Ala        | gga<br>Gly<br>560 | 1680 |
| ttt<br>Phe | gtt<br>Val        | gcc<br>Ala        | agc<br>Ser        | atc<br>Ile<br>565 | aat<br>Asn        | gaa<br>Glu        | Gly<br>333        | atg<br>Met        | acc<br>Thr<br>570 | cgc<br>Arg        | tgg<br>Trp        | ttc<br>Phe        | tca<br>Ser        | cgc<br>Arg<br>575 | tgc<br>Cys        | 1728 |
| ata<br>Ile | ttt<br>Phe        | cag<br>Gln        | gat<br>Asp<br>580 | aga<br>Arg        | gga<br>Gly        | cag<br>Gln        | gag<br>Glu        | ctg<br>Leu<br>585 | gta<br>Val        | gat<br>Asp        | Gly<br>aaa        | ctc<br>Leu        | aaa<br>Lys<br>590 | gtc<br>Val        | tgc<br>Cys        | 1776 |
| ctg<br>Leu | caa<br>Gln        | gcg<br>Ala<br>595 | gct<br>Ala        | ctg<br>Leu        | agg<br>Arg        | gct<br>Ala        | tgg<br>Trp<br>600 | aat<br>Asn        | agc<br>Ser        | tgc<br>Cys        | aat<br>Asn        | gag<br>Glu<br>605 | tac<br>Tyr        | atg<br>Met        | ccc<br>Pro        | 1824 |
| agc<br>Ser | cgg<br>Arg<br>610 | atc<br>Ile        | atc<br>Ile        | gtg<br>Val        | tac<br>Tyr        | cgc<br>Arg<br>615 | gat<br>Asp        | ggc               | gta<br>Val        | gga<br>Gly        | gac<br>Asp<br>620 | ggc<br>Gly        | cag<br>Gln        | ctg<br>Leu        | aaa<br>Lys        | 1872 |
| aca<br>Thr | ctg<br>Leu        | gtg<br>Val        | aac<br>Asn        | tac<br>Tyr        | gaa<br>Glu        | gtg<br>Val        | cca<br>Pro        | cag<br>Gln        | ttt<br>Phe        | ttg<br>Leu        | gat<br>Asp        | tgt<br>Cys        | cta<br>Leu        | aaa<br>Lys        | tcc<br>Ser        | 1920 |

| 625   | 630                                     |             | 635                                       | 640  |
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| Arg Val Asn Th                                    |   | -           | gga gga aga ctt<br>Gly Gly Arg Leu<br>670 |      |
|   |   |             | gtt acc aga cca<br>Val Thr Arg Pro<br>685 |      |
|   |   | Gln Ala Val | aga agt ggt agt<br>Arg Ser Gly Ser<br>700 |      |
| 5 9   | _                                       | _           | agc ggc ctg aag<br>Ser Gly Leu Lys<br>715 | -    |
| . "   | _                                       |             | cac atc tat tac<br>His Ile Tyr Tyr        |      |
| Pro Gly Val I                                     |   | _           | cag tac gcc cac<br>Gln Tyr Ala His<br>750 |      |
| 5 1 2   |   |             | gag cca aat ctg<br>Glu Pro Asn Leu<br>765 | _    |
|   | tt tac tac cto<br>eu Tyr Tyr Lei<br>775 | 1           |   | 2328 |
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85 90 95 Ser Lys Thr Arg Asn Gly Glu Asp Val Arg Ile Thr Ile Thr Leu Thr i. Asn Glu Leu Pro Pro Thr Ser Pro Thr Cys Leu Gln Phe Tyr Asn Ile Ile Phe Arg Arg Leu Leu Lys Ile Met Asn Leu Gln Gln Ile Gly Arg Asn Tyr Tyr Asn Pro Asn Asp Pro Ile Asp Ile Pro Ser His Arg Leu Val Ile Trp Pro Gly Phe Thr Thr Ser Ile Leu Gln Tyr Glu Asn Ser Ile Met Leu Cys Thr Asp Val Ser His Lys Val Leu Arg Ser Glu Thr 

Val Leu Asp Phe Met Phe Asn Phe Tyr His Gln Thr Glu Glu His Lys Phe Gln Glu Gln Val Ser Lys Glu Leu Ile Gly Leu Val Val Leu Thr Lys Tyr Asn Asn Lys Thr Tyr Arg Val Asp Asp Ile Asp Trp Asp Gln Asn Pro Lys Ser Thr Phe Lys Lys Ala Asp Gly Ser Glu Val Ser Phe Leu Glu Tyr Tyr Arg Lys Gln Tyr Asn Gln Glu Ile Thr Asp Leu Lys 🗐n Pro Val Leu Val Ser Gln Pro Lys Arg Arg Arg Gly Pro Gly Gly Thr Leu Pro Gly Pro Ala Met Leu Ile Pro Glu Leu Cys Tyr Leu Thr Taly Leu Thr Asp Lys Met Arg Asn Asp Phe Asn Val Met Lys Asp Leu Ala Val His Thr Arg Leu Thr Pro Glu Gln Arg Gln Arg Glu Val Gly Arg Leu Ile Asp Tyr Ile His Lys Asn Asp Asn Val Gln Arg Glu Leu Arg Asp Trp Gly Leu Ser Phe Asp Ser Asn Leu Leu Ser Phe Ser Gly 

Arg Ile Leu Gln Thr Glu Lys Ile His Gln Gly Gly Lys Thr Phe Asp Tyr Asn Pro Gln Phe Ala Asp Trp Ser Lys Glu Thr Arg Gly Ala Pro Leu Ile Ser Val Lys Pro Leu Asp Asn Trp Leu Leu Ile Tyr Thr Arg Arg Asn Tyr Glu Ala Ala Asn Ser Leu Ile Gln Asn Leu Phe Lys Val Thr Pro Ala Met Gly Met Gln Met Arg Lys Ala Ile Met Ile Glu Val Asp Asp Arg Thr Glu Ala Tyr Leu Arg Val Leu Gln Gln Lys Val Thr 索la Asp Thr Gln Ile Val Val Cys Leu Leu Ser Ser Asn Arg Lys Asp H Lys Tyr Asp Ala Ile Lys Lys Tyr Leu Cys Thr Asp Cys Pro Thr Pro
485 490 495 N Ser Gln Cys Val Val Ala Arg Thr Leu Gly Lys Gln Gln Thr Val Met Ala Ile Ala Thr Lys Ile Ala Leu Gln Met Asn Cys Lys Met Gly Gly Glu Leu Trp Arg Val Asp Ile Pro Leu Lys Leu Val Met Ile Val Gly Ile Asp Cys Tyr His Asp Met Thr Ala Gly Arg Arg Ser Ile Ala Gly 

Phe Val Ala Ser Ile Asn Glu Gly Met Thr Arg Trp Phe Ser Arg Cys Ile Phe Gln Asp Arg Gly Gln Glu Leu Val Asp Gly Leu Lys Val Cys Leu Gln Ala Ala Leu Arg Ala Trp Asn Ser Cys Asn Glu Tyr Met Pro Ser Arg Ile Ile Val Tyr Arg Asp Gly Val Gly Asp Gly Gln Leu Lys Thr Leu Val Asn Tyr Glu Val Pro Gln Phe Leu Asp Cys Leu Lys Ser 625 630 635 640 The Gly Arg Gly Tyr Asn Pro Arg Leu Thr Val Ile Val Val Lys Lys Arg Val Asn Thr Arg Phe Phe Ala Gln Ser Gly Gly Arg Leu Gln Asn Pro Leu Pro Gly Thr Val Ile Asp Val Glu Val Thr Arg Pro Glu Trp Tyr Asp Phe Phe Ile Val Ser Gln Ala Val Arg Ser Gly Ser Val Ser Pro Thr His Tyr Asn Val Ile Tyr Asp Asn Ser Gly Leu Lys Pro Asp His Ile Gln Arg Leu Thr Tyr Lys Leu Cys His Ile Tyr Tyr Asn Trp 

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Pro Gly Val Ile Arg Val Pro Ala Pro Cys Gln Tyr Ala His Lys Leu
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                                  745
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Gln Gly Thr Val Ser Pro Thr Ser Tyr Asn Val Leu Tyr Ser Ser Met 770 775 780

Cly Leu Ser Pro Glu Lys Met Gln Lys Leu Thr Tyr Lys Met Cys His 795 795 800

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